Andrew D Ellington

List of Publications by Year in Descending Order

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The third column is the impact factor (IF) of the journal, and the fourth column is the number of citations of the article.

299	26,597	74	159
papers	citations	h-index	g-index
326 ext. papers	29,771 ext. citations	11.5 avg, IF	7.31 L-index

#	Paper	IF	Citations
299	Evolving a Generalist Biosensor for Bicyclic Monoterpenes ACS Synthetic Biology, 2022,	5.7	2
298	Making Security Viral: Shifting Engineering Biology Culture and Publishing <i>ACS Synthetic Biology</i> , 2022 , 11, 522-527	5.7	О
297	Preparation and Use of Cellular Reagents: A Low-resource Molecular Biology Reagent Platform <i>Current Protocols</i> , 2022 , 2, e387		O
296	Machine learning-aided engineering of hydrolases for PET depolymerization <i>Nature</i> , 2022 , 604, 662-66	67 50.4	23
295	Developing predictive hybridization models for phosphorothioate oligonucleotides using high-resolution melting <i>PLoS ONE</i> , 2022 , 17, e0268575	3.7	
294	Learning the local landscape of protein structures with convolutional neural networks. <i>Journal of Biological Physics</i> , 2021 , 47, 435-454	1.6	1
293	Improved Bst DNA Polymerase Variants Derived a Machine Learning Approach. <i>Biochemistry</i> , 2021 ,	3.2	3
292	Heat adaptation of phage T7 under an extended genetic code Virus Evolution, 2021, 7, veab100	3.7	0
291	Synthetic repertoires derived from convalescent COVID-19 patients enable discovery of SARS-CoV-2 neutralizing antibodies and a novel quaternary binding modality 2021 ,		4
290	High-Surety Isothermal Amplification and Detection of SARS-CoV-2. MSphere, 2021, 6,	5	22
289	Guiding Ethical Principles in Engineering Biology Research. ACS Synthetic Biology, 2021 , 10, 907-910	5.7	4
288	Directed Evolution of an Improved Aminoacyl-tRNA Synthetase for Incorporation of L-3,4-Dihydroxyphenylalanine (L-DOPA). <i>Angewandte Chemie - International Edition</i> , 2021 , 60, 14811-14	816.4	4
287	Directed Evolution of an Improved Aminoacyl-tRNA Synthetase for Incorporation of L-3,4-Dihydroxyphenylalanine (L-DOPA). <i>Angewandte Chemie</i> , 2021 , 133, 14937-14942	3.6	4
286	Minimizing Leakage in Stacked Strand Exchange Amplification Circuits. <i>ACS Synthetic Biology</i> , 2021 , 10, 1277-1283	5.7	0
285	Producing molecular biology reagents without purification. <i>PLoS ONE</i> , 2021 , 16, e0252507	3.7	2
284	Ribosome-mediated incorporation of fluorescent amino acids into peptides in vitro. <i>Chemical Communications</i> , 2021 , 57, 2661-2664	5.8	5
283	Recombineering and MAGE Nature Reviews Methods Primers, 2021, 1,		13

282	Delineation of the Ancestral Tus-Dependent Replication Fork Trap <i>International Journal of Molecular Sciences</i> , 2021 , 22,	6.3	1	
281	Dynamic Programming of a DNA Walker Controlled by Protons. <i>ACS Nano</i> , 2020 , 14, 4007-4013	16.7	36	
280	Site-specific 5-hydroxytryptophan incorporation into apolipoprotein A-I impairs cholesterol efflux activity and high-density lipoprotein biogenesis. <i>Journal of Biological Chemistry</i> , 2020 , 295, 4836-4848	5.4	8	
279	Engineered symbionts activate honey bee immunity and limit pathogens. <i>Science</i> , 2020 , 367, 573-576	33.3	81	
278	A facile technology for the high-throughput sequencing of the paired VH:VL and TCRUTCRH repertoires. <i>Science Advances</i> , 2020 , 6, eaay9093	14.3	12	
277	Ribosomal incorporation of cyclic Elamino acids into peptides using in vitro translation. <i>Chemical Communications</i> , 2020 , 56, 5597-5600	5.8	13	
276	Emulsion-based directed evolution of enzymes and proteins in yeast. <i>Methods in Enzymology</i> , 2020 , 643, 87-110	1.7		
275	Oligonucleotide-functionalized hydrogels for sustained release of small molecule (aptamer) therapeutics. <i>Acta Biomaterialia</i> , 2020 , 102, 315-325	10.8	12	
274	Discovery of Novel Gain-of-Function Mutations Guided by Structure-Based Deep Learning. <i>ACS Synthetic Biology</i> , 2020 , 9, 2927-2935	5.7	18	
273	One-Enzyme Reverse Transcription qPCR Using Taq DNA Polymerase. <i>Biochemistry</i> , 2020 , 59, 4638-464	53.2	9	
272	How a B family DNA polymerase has been evolved to copy RNA. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2020 , 117, 21274-21280	11.5	3	
271	Bringing Microscopy-By-Sequencing into View. <i>Trends in Biotechnology</i> , 2020 , 38, 154-162	15.1	2	
270	Employing 25-Residue Docking Motifs from Modular Polyketide Synthases as Orthogonal Protein Connectors. <i>ACS Synthetic Biology</i> , 2019 , 8, 2017-2024	5.7	3	
269	Single-Molecule Mechanistic Study of Enzyme Hysteresis. ACS Central Science, 2019, 5, 1691-1698	16.8	11	
268	Synthetic evolution. <i>Nature Biotechnology</i> , 2019 , 37, 730-743	44.5	36	
267	Pattern Generation with Nucleic Acid Chemical Reaction Networks. <i>Chemical Reviews</i> , 2019 , 119, 6370-	6 383 1	32	
266	Aptamers in Education: Undergraduates Make Aptamers and Acquire 21st Century Skills Along the Way. <i>Sensors</i> , 2019 , 19,	3.8	1	
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264	Expanding the limits of the second genetic code with ribozymes. <i>Nature Communications</i> , 2019 , 10, 509	7 17.4	54
263	Retrons and their applications in genome engineering. <i>Nucleic Acids Research</i> , 2019 , 47, 11007-11019	20.1	35
262	Synthetic GPCRs and signal transduction cascades. <i>Emerging Topics in Life Sciences</i> , 2019 , 3, 609-614	3.5	O
261	Hachimoji DNA and RNA: A genetic system with eight building blocks. <i>Science</i> , 2019 , 363, 884-887	33.3	193
2 60	Reprogramming the brain with synthetic neurobiology. Current Opinion in Biotechnology, 2019, 58, 37-4	411.4	2
259	Supercharging enables organized assembly of synthetic biomolecules. <i>Nature Chemistry</i> , 2019 , 11, 204-	2 17 .6	48
258	Synthesis of Ferrocene Derivatives Allowing Linear Free Energy Studies of Redox Potentials. <i>Helvetica Chimica Acta</i> , 2019 , 102, e1800186	2	2
257	Evolving Bacterial Fitness with an Expanded Genetic Code. <i>Scientific Reports</i> , 2018 , 8, 3288	4.9	4
256	Directed evolution of a synthetic phylogeny of programmable Trp repressors. <i>Nature Chemical Biology</i> , 2018 , 14, 361-367	11.7	36
255	Construction of synthetic T7 RNA polymerase expression systems. <i>Methods</i> , 2018 , 143, 110-120	4.6	11
254	Strand Displacement Probes Combined with Isothermal Nucleic Acid Amplification for Instrument-Free Detection from Complex Samples. <i>Analytical Chemistry</i> , 2018 , 90, 6580-6586	7.8	66
253	Evolution of a Thermophilic Strand-Displacing Polymerase Using High-Temperature Isothermal Compartmentalized Self-Replication. <i>Biochemistry</i> , 2018 , 57, 4607-4619	3.2	16
252	Continuous directed evolution for strain and protein engineering. <i>Current Opinion in Biotechnology</i> , 2018 , 53, 158-163	11.4	25
251	Portable platform for rapid in-field identification of human fecal pollution in water. <i>Water Research</i> , 2018 , 131, 186-195	12.5	22
250	Functional interrogation and mining of natively paired human V:V antibody repertoires. <i>Nature Biotechnology</i> , 2018 , 36, 152-155	44.5	80
249	Genetic Engineering of Bee Gut Microbiome Bacteria with a Toolkit for Modular Assembly of Broad-Host-Range Plasmids. <i>ACS Synthetic Biology</i> , 2018 , 7, 1279-1290	5.7	46
248	Fingerprinting Non-Terran Biosignatures. <i>Astrobiology</i> , 2018 , 18, 915-922	3.7	24
247	In Vitro Transcription Networks Based on Hairpin Promoter Switches. <i>ACS Synthetic Biology</i> , 2018 , 7, 1937-1945	5.7	11

246	Cellular reagents for diagnostics and synthetic biology. <i>PLoS ONE</i> , 2018 , 13, e0201681	3.7	13
245	Custom selenoprotein production enabled by laboratory evolution of recoded bacterial strains. <i>Nature Biotechnology</i> , 2018 , 36, 624-631	44.5	21
244	Compartmentalized Self-Replication for Evolution of a DNA Polymerase. <i>Current Protocols in Chemical Biology</i> , 2018 , 10, 1-17	1.8	3
243	How to Balance the Many Roles of tRNAs During the Creation of New Genetic Codes. <i>FASEB Journal</i> , 2018 , 32, 105.2	0.9	
242	A highly parallel strategy for storage of digital information in living cells. <i>BMC Biotechnology</i> , 2018 , 18, 64	3.5	7
241	Simultaneous Detection of Different Zika Virus Lineages via Molecular Computation in a Point-of-Care Assay. <i>Viruses</i> , 2018 , 10,	6.2	10
240	Effective design principles for leakless strand displacement systems. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2018 , 115, E12182-E12191	11.5	59
239	Viral attenuation by engineered protein fragmentation. Virus Evolution, 2018, 4, vey017	3.7	1
238	Retroelement-Based Genome Editing and Evolution. ACS Synthetic Biology, 2018, 7, 2600-2611	5.7	27
237	Predicting Evolution of the Transcription Regulatory Network in a Bacteriophage. <i>Genome Biology and Evolution</i> , 2018 , 10, 2614-2628	3.9	
236	Direct nucleic acid analysis of mosquitoes for high fidelity species identification and detection of Wolbachia using a cellphone. <i>PLoS Neglected Tropical Diseases</i> , 2018 , 12, e0006671	4.8	15
235	Phosphorothioated Primers Lead to Loop-Mediated Isothermal Amplification at Low Temperatures. <i>Analytical Chemistry</i> , 2018 , 90, 8290-8294	7.8	39
234	Amplicon Competition Enables End-Point Quantitation of Nucleic Acids Following Isothermal Amplification. <i>ChemBioChem</i> , 2017 , 18, 1692-1695	3.8	13
233	Characterization of trimethoprim resistant dihydrofolate reductase mutants by mass spectrometry and inhibition by propargyl-linked antifolates. <i>Chemical Science</i> , 2017 , 8, 4062-4072	9.4	24
232	Synthetic DNA Synthesis and Assembly: Putting the Synthetic in Synthetic Biology. <i>Cold Spring Harbor Perspectives in Biology</i> , 2017 , 9,	10.2	166
231	Coupling Sensitive Nucleic Acid Amplification with Commercial Pregnancy Test Strips. <i>Angewandte Chemie</i> , 2017 , 129, 1012-1016	3.6	16
230	Coupling Sensitive Nucleic Acid Amplification with Commercial Pregnancy Test Strips. <i>Angewandte Chemie - International Edition</i> , 2017 , 56, 992-996	16.4	98
229	Genetic alphabet expansion transcription generating functional RNA molecules containing a five-letter alphabet including modified unnatural and natural base nucleotides by thermostable T7 RNA polymerase variants. <i>Chemical Communications</i> , 2017 , 53, 12309-12312	5.8	15

228	Purification of single-stranded DNA by co-polymerization with acrylamide and electrophoresis. <i>BioTechniques</i> , 2017 , 62, 275-282	2.5	13
227	The Design Space of Strand Displacement Cascades with Toehold-Size Clamps. <i>Lecture Notes in Computer Science</i> , 2017 , 64-81	0.9	5
226	Charge Shielding Prevents Aggregation of Supercharged GFP Variants at High Protein Concentration. <i>Molecular Pharmaceutics</i> , 2017 , 14, 3269-3280	5.6	17
225	A Simple, Cleated DNA Walker That Hangs on to Surfaces. ACS Nano, 2017, 11, 8047-8054	16.7	82
224	Compartmentalized partnered replication for the directed evolution of genetic parts and circuits. <i>Nature Protocols</i> , 2017 , 12, 2493-2512	18.8	18
223	Differential array sensing for cancer cell classification and novelty detection. <i>Organic and Biomolecular Chemistry</i> , 2017 , 15, 9866-9874	3.9	16
222	Massively Parallel Biophysical Analysis of CRISPR-Cas Complexes on Next Generation Sequencing Chips. <i>Cell</i> , 2017 , 170, 35-47.e13	56.2	62
221	Evolving Orthogonal Suppressor tRNAs To Incorporate Modified Amino Acids. <i>ACS Synthetic Biology</i> , 2017 , 6, 108-119	5.7	22
220	Synthesis of alanyl nucleobase amino acids and their incorporation into proteins. <i>Bioorganic and Medicinal Chemistry</i> , 2016 , 24, 4177-4187	3.4	6
219	Design and engineering of a transmissible antiviral defense. <i>Journal of Biological Engineering</i> , 2016 , 10, 12	6.3	
218	Molecular-level analysis of the serum antibody repertoire in young adults before and after seasonal influenza vaccination. <i>Nature Medicine</i> , 2016 , 22, 1456-1464	50.5	186
217	Synthesis and structural analyses of phenylethynyl-substituted tris(2-pyridylmethyl)amines and their copper(ii) complexes. <i>Dalton Transactions</i> , 2016 , 45, 10585-98	4.3	3
216	Synthetic evolutionary origin of a proofreading reverse transcriptase. <i>Science</i> , 2016 , 352, 1590-3	33.3	84
215	Engineering Signaling Aptamers That Rely on Kinetic Rather Than Equilibrium Competition. <i>Analytical Chemistry</i> , 2016 , 88, 2250-7	7.8	14
214	Addicting diverse bacteria to a noncanonical amino acid. <i>Nature Chemical Biology</i> , 2016 , 12, 138-40	11.7	37
213	Ultra-high-throughput sequencing of the immune receptor repertoire from millions of lymphocytes. <i>Nature Protocols</i> , 2016 , 11, 429-42	18.8	103
212	An in vitro selection for small molecule induced switching RNA molecules. <i>Methods</i> , 2016 , 106, 51-7	4.6	7
211	Virus wars: using one virus to block the spread of another. <i>PeerJ</i> , 2016 , 4, e2166	3.1	3

210	Recent advances in synthetic biosafety. F1000Research, 2016, 5,	3.6	14
209	Strand-Exchange Nucleic Acid Circuitry with Enhanced Thermo-and Structure- Buffering Abilities Turns Gene Diagnostics Ultra-Reliable and Environmental Compatible. <i>Scientific Reports</i> , 2016 , 6, 36605	4.9	13
208	A primerless molecular diagnostic: phosphorothioated-terminal hairpin formation and self-priming extension (PS-THSP). <i>Analytical and Bioanalytical Chemistry</i> , 2016 , 408, 8583-8591	4.4	11
207	Large-scale sequence and structural comparisons of human naive and antigen-experienced antibody repertoires. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2016 , 113, E2636-45	11.5	118
206	Expanded Genetic Codes Create New Mutational Routes to Rifampicin Resistance in Escherichia coli. <i>Molecular Biology and Evolution</i> , 2016 , 33, 2054-63	8.3	9
205	Design, synthesis, and application of Spinach molecular beacons triggered by strand displacement. <i>Methods in Enzymology</i> , 2015 , 550, 215-49	1.7	6
204	Synthetic biology: Six pack and stack. <i>Nature Chemistry</i> , 2015 , 7, 617-9	17.6	2
203	A Sweet Spot for Molecular Diagnostics: Coupling Isothermal Amplification and Strand Exchange Circuits to Glucometers. <i>Scientific Reports</i> , 2015 , 5, 11039	4.9	54
202	Structural Characterization of Dihydrofolate Reductase Complexes by Top-Down Ultraviolet Photodissociation Mass Spectrometry. <i>Journal of the American Chemical Society</i> , 2015 , 137, 9128-35	16.4	54
201	Robust strand exchange reactions for the sequence-specific, real-time detection of nucleic acid amplicons. <i>Analytical Chemistry</i> , 2015 , 87, 3314-20	7.8	96
200	A microbial model of economic trading and comparative advantage. <i>Journal of Theoretical Biology</i> , 2015 , 364, 326-43	2.3	11
199	In Vitro Selection for Small-Molecule-Triggered Strand Displacement and Riboswitch Activity. <i>ACS Synthetic Biology</i> , 2015 , 4, 1144-50	5.7	19
198	High-affinity RNA Aptamers Against the HIV-1 Protease Inhibit Both In Vitro Protease Activity and Late Events of Viral Replication. <i>Molecular Therapy - Nucleic Acids</i> , 2015 , 4, e228	10.7	27
197	Transcription yield of fully 2Fmodified RNA can be increased by the addition of thermostabilizing mutations to T7 RNA polymerase mutants. <i>Nucleic Acids Research</i> , 2015 , 43, 7480-8	20.1	44
196	RNA as a conception. <i>Rna</i> , 2015 , 21, 608	5.8	
195	In-depth determination and analysis of the human paired heavy- and light-chain antibody repertoire. <i>Nature Medicine</i> , 2015 , 21, 86-91	50.5	259
194	Directed Evolution of a Panel of Orthogonal T7 RNA Polymerase Variants for in Vivo or in Vitro Synthetic Circuitry. <i>ACS Synthetic Biology</i> , 2015 , 4, 1070-6	5.7	36
193	Next-generation sequencing as input for chemometrics in differential sensing routines. Angewandte Chemie - International Edition, 2015, 54, 6339-42	16.4	5

192	Next-Generation Sequencing as Input for Chemometrics in Differential Sensing Routines. <i>Angewandte Chemie</i> , 2015 , 127, 6437-6440	3.6	3
191	Real-time sequence-validated loop-mediated isothermal amplification assays for detection of Middle East respiratory syndrome coronavirus (MERS-CoV). <i>PLoS ONE</i> , 2015 , 10, e0123126	3.7	96
190	One-step tumor detection from dynamic morphology tracking on aptamer-grafted surfaces. <i>Technology</i> , 2015 , 3, 194-200	3	6
189	Industrialization of Biology. ACS Synthetic Biology, 2015 , 4, 1053-5	5.7	12
188	Landscape-Based Biology. <i>Journal of Molecular Evolution</i> , 2015 , 81, 144-5	3.1	1
187	Chemical Tools To Decipher Regulation of Phosphatases by Proline Isomerization on Eukaryotic RNA Polymerase II. <i>ACS Chemical Biology</i> , 2015 , 10, 2405-14	4.9	18
186	3D Printing with Nucleic Acid Adhesives. ACS Biomaterials Science and Engineering, 2015, 1, 19-26	5.5	21
185	Controlled assembly of artificial protein-protein complexes via DNA duplex formation. <i>Bioconjugate Chemistry</i> , 2015 , 26, 427-34	6.3	1
184	Fine-tuning citrate synthase flux potentiates and refines metabolic innovation in the Lenski evolution experiment. <i>ELife</i> , 2015 , 4,	8.9	48
183	Alternative ELISA Using a RNA Aptamer against Calf Intestinal Alkaline Phosphatase. <i>FASEB Journal</i> , 2015 , 29, 562.6	0.9	
182	Diagnostic applications of nucleic acid circuits. Accounts of Chemical Research, 2014, 47, 1825-35	24.3	225
181	Structure-based non-canonical amino acid design to covalently crosslink an antibody-antigen complex. <i>Journal of Structural Biology</i> , 2014 , 185, 215-22	3.4	18
180	Directed evolution of genetic parts and circuits by compartmentalized partnered replication. <i>Nature Biotechnology</i> , 2014 , 32, 97-101	44.5	103
179	Bacteriophages use an expanded genetic code on evolutionary paths to higher fitness. <i>Nature Chemical Biology</i> , 2014 , 10, 178-80	11.7	39
178	A proteomic survey of widespread protein aggregation in yeast. <i>Molecular BioSystems</i> , 2014 , 10, 851-86	51	40
177	A Spinach molecular beacon triggered by strand displacement. <i>Rna</i> , 2014 , 20, 1183-94	5.8	42
176	Progress Report on the Generation of Polyfunctional Microscale Particles for Programmed Self-Assembly. <i>Chemistry of Materials</i> , 2014 , 26, 1457-1462	9.6	4
175	Directed evolution of the substrate specificity of biotin ligase. <i>Biotechnology and Bioengineering</i> , 2014 , 111, 1071-81	4.9	11

(2013-2014)

174	Mismatches Improve the Performance of Strand-Displacement Nucleic Acid Circuits. <i>Angewandte Chemie</i> , 2014 , 126, 1876-1879	3.6	50
173	Modeling Scalable Pattern Generation in DNA Reaction Networks. <i>Natural Computing</i> , 2014 , 13, 583-59	5 1.3	4
172	In vitro selection using modified or unnatural nucleotides. <i>Current Protocols in Nucleic Acid Chemistry</i> , 2014 , 56, 9.6.1-33	0.5	22
171	Design and selection of a synthetic operon. ACS Synthetic Biology, 2014, 3, 410-5	5.7	2
170	Library generation by gene shuffling. Current Protocols in Molecular Biology, 2014, 105, Unit 15.12.	2.9	12
169	Recursive genomewide recombination and sequencing reveals a key refinement step in the evolution of a metabolic innovation in Escherichia coli. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2014 , 111, 2217-22	11.5	50
168	An amino acid depleted cell-free protein synthesis system for the incorporation of non-canonical amino acid analogs into proteins. <i>Journal of Biotechnology</i> , 2014 , 178, 12-22	3.7	23
167	Differential Sensing of MAP Kinases Using SOX-Peptides. <i>Angewandte Chemie</i> , 2014 , 126, 14288-14292	3.6	5
166	Mismatches improve the performance of strand-displacement nucleic Acid circuits. <i>Angewandte Chemie - International Edition</i> , 2014 , 53, 1845-8	16.4	120
165	Design and application of cotranscriptional non-enzymatic RNA circuits and signal transducers. <i>Nucleic Acids Research</i> , 2014 , 42, e58	20.1	61
164	Exquisite allele discrimination by toehold hairpin primers. <i>Nucleic Acids Research</i> , 2014 , 42, e120	20.1	5
163	Analyzing machupo virus-receptor binding by molecular dynamics simulations. <i>PeerJ</i> , 2014 , 2, e266	3.1	7
162	Proliferation and migration of tumor cells in tapered channels. <i>Biomedical Microdevices</i> , 2013 , 15, 635-6	5 43 7	26
161	Generalized bacterial genome editing using mobile group II introns and Cre-lox. <i>Molecular Systems Biology</i> , 2013 , 9, 685	12.2	63
160	Pattern transformation with DNA circuits. <i>Nature Chemistry</i> , 2013 , 5, 1000-5	17.6	102
159	In vitro selection of proteins via emulsion compartments. <i>Methods</i> , 2013 , 60, 75-80	4.6	22
158	Real-time detection of isothermal amplification reactions with thermostable catalytic hairpin assembly. <i>Journal of the American Chemical Society</i> , 2013 , 135, 7430-3	16.4	215
157	Effect of Complementary Nucleobase Interactions on the Copolymer Composition of RAFT Copolymerizations <i>ACS Macro Letters</i> , 2013 , 2, 581-586	6.6	54

156	DNA detection using origami paper analytical devices. <i>Analytical Chemistry</i> , 2013 , 85, 9713-20	7.8	102
155	Continuous in vitro evolution of a ribozyme ligase: a model experiment for the evolution of a biomolecule. <i>Biochemistry and Molecular Biology Education</i> , 2013 , 41, 433-42	1.3	
154	Stacking nonenzymatic circuits for high signal gain. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2013 , 110, 5386-91	11.5	182
153	Alternative computational protocols for supercharging protein surfaces for reversible unfolding and retention of stability. <i>PLoS ONE</i> , 2013 , 8, e64363	3.7	50
152	A general RNA motif for cellular transfection. <i>Molecular Therapy</i> , 2012 , 20, 616-24	11.7	29
151	An in vitro autogene. ACS Synthetic Biology, 2012 , 1, 190-6	5.7	14
150	Probing spatial organization of DNA strands using enzyme-free hairpin assembly circuits. <i>Journal of the American Chemical Society</i> , 2012 , 134, 13918-21	16.4	189
149	Origins for Everyone. <i>Evolution: Education and Outreach</i> , 2012 , 5, 361-366	1.6	6
148	Exploration of plasticizer and plastic explosive detection and differentiation with serum albumin cross-reactive arrays. <i>Chemical Science</i> , 2012 , 3, 1773	9.4	24
147	Adapting enzyme-free DNA circuits to the detection of loop-mediated isothermal amplification reactions. <i>Analytical Chemistry</i> , 2012 , 84, 8371-7	7.8	78
146	Spatial control of DNA reaction networks by DNA sequence. <i>Molecules</i> , 2012 , 17, 13390-402	4.8	13
145	DNA circuits as amplifiers for the detection of nucleic acids on a paperfluidic platform. <i>Lab on A Chip</i> , 2012 , 12, 2951-8	7.2	74
144	A fully-electronic charge-based DNA sequencing CMOS biochip 2012 ,		13
143	Structure-based design of supercharged, highly thermoresistant antibodies. <i>Chemistry and Biology</i> , 2012 , 19, 449-55		108
142	Ribozymes as Molecular Biology Reagents 2012 , 293-312		
141	Rational, modular adaptation of enzyme-free DNA circuits to multiple detection methods. <i>Nucleic Acids Research</i> , 2011 , 39, e110	20.1	363
140	Inhibition of cell proliferation by an anti-EGFR aptamer. PLoS ONE, 2011, 6, e20299	3.7	132
139	Identifying protein variants with cross-reactive aptamer arrays. ChemBioChem, 2011, 12, 2021-4	3.8	15

138	Aptamers as therapeutics. <i>Nature Reviews Drug Discovery</i> , 2010 , 9, 537-50	64.1	1428
137	Aptamer antagonists of myelin-derived inhibitors promote axon growth. <i>PLoS ONE</i> , 2010 , 5, e9726	3.7	10
136	In vitro selection of RNA aptamers to a protein target by filter immobilization. <i>Current Protocols in Nucleic Acid Chemistry</i> , 2010 , Chapter 9, Unit 9.3.1-27	0.5	7
135	In vitro selection of RNA aptamers to a small molecule target. <i>Current Protocols in Nucleic Acid Chemistry</i> , 2010 , Chapter 9, Unit 9.5.1-23	0.5	7
134	Shaping up nucleic acid computation. <i>Current Opinion in Biotechnology</i> , 2010 , 21, 392-400	11.4	48
133	Beyond allostery: Catalytic regulation of a deoxyribozyme through an entropy-driven DNA amplifier. <i>Journal of Systems Chemistry</i> , 2010 , 1, 13		31
132	In vitro selection of RNA aptamers to a protein target by filter immobilization. <i>Current Protocols in Molecular Biology</i> , 2009 , Chapter 24, Unit 24.3	2.9	5
131	Directed evolution of proteins in vitro using compartmentalization in emulsions. <i>Current Protocols in Molecular Biology</i> , 2009 , Chapter 24, Unit 24.6	2.9	8
130	Design principles for ligand-sensing, conformation-switching ribozymes. <i>PLoS Computational Biology</i> , 2009 , 5, e1000620	5	42
129	Modelling amorphous computations with transcription networks. <i>Journal of the Royal Society Interface</i> , 2009 , 6 Suppl 4, S523-33	4.1	18
128	Kinetic optimization of a protein-responsive aptamer beacon. <i>Biotechnology and Bioengineering</i> , 2009 , 103, 1049-59	4.9	21
127	Direct selection for ribozyme cleavage activity in cells. <i>Rna</i> , 2009 , 15, 2035-45	5.8	20
126	Evolutionary origins and directed evolution of RNA. <i>International Journal of Biochemistry and Cell Biology</i> , 2009 , 41, 254-65	5.6	59
125	A synthetic genetic edge detection program. <i>Cell</i> , 2009 , 137, 1272-81	56.2	37 ²
124	Applications of aptamers as sensors. Annual Review of Analytical Chemistry, 2009, 2, 241-64	12.5	633
123	Technical and biological issues relevant to cell typing with aptamers. <i>Journal of Proteome Research</i> , 2009 , 8, 2438-48	5.6	79
122	Using RNA aptamers and the proximity ligation assay for the detection of cell surface antigens. <i>Methods in Molecular Biology</i> , 2009 , 504, 385-98	1.4	17
121	Aptamer Microarrays 2009 , 287-308		

120	Aptamer-targeted gold nanoparticles as molecular-specific contrast agents for reflectance imaging. <i>Bioconjugate Chemistry</i> , 2008 , 19, 1309-12	6.3	140
119	Aptamer amplification: divide and signal. Expert Opinion on Medical Diagnostics, 2008, 2, 1333-46		7
118	Man versus machine versus ribozyme. <i>PLoS Biology</i> , 2008 , 6, e132	9.7	1
117	NUCLEIC ACIDS FOR REAGENTLESS BIOSENSORS 2008 , 493-541		2
116	Selection of fluorescent aptamer beacons that light up in the presence of zinc. <i>Analytical and Bioanalytical Chemistry</i> , 2008 , 390, 1067-75	4.4	120
115	Bioinformatic analysis of the contribution of primer sequences to aptamer structures. <i>Journal of Molecular Evolution</i> , 2008 , 67, 95-102	3.1	62
114	Real-time PCR detection of protein analytes with conformation-switching aptamers. <i>Analytical Biochemistry</i> , 2008 , 380, 164-73	3.1	40
113	Directed evolution of streptavidin variants using in vitro compartmentalization. <i>Chemistry and Biology</i> , 2008 , 15, 979-89		22
112	What's so great about RNA?. ACS Chemical Biology, 2007, 2, 445-8	4.9	7
111	Real-time rolling circle amplification for protein detection. <i>Analytical Chemistry</i> , 2007 , 79, 3320-9	7.8	166
110	Ribozyme catalysis of metabolism in the RNA world. Chemistry and Biodiversity, 2007, 4, 633-55	2.5	138
109	Synthetic RNA circuits. <i>Nature Chemical Biology</i> , 2007 , 3, 23-8	11.7	43
108	Computational selection of nucleic acid biosensors via a slip structure model. <i>Biosensors and Bioelectronics</i> , 2007 , 22, 1939-47	11.8	40
107	Deoxyribozymes that recode sequence information. <i>Nucleic Acids Research</i> , 2006 , 34, 2166-72	20.1	16
106	Aptazymes: Allosteric Ribozymes and Deoxyribozymes as Biosensors 2006 , 290-310		6
105	Aptamer:toxin conjugates that specifically target prostate tumor cells. <i>Cancer Research</i> , 2006 , 66, 5989) -92 .1	240
104	Ribozyme-mediated signal augmentation on a mass-sensitive biosensor. <i>Journal of the American Chemical Society</i> , 2006 , 128, 15936-7	16.4	40
103	Labeling tumor cells with fluorescent nanocrystal-aptamer bioconjugates. <i>Biosensors and Bioelectronics</i> , 2006 , 21, 1859-66	11.8	131

(2004-2006)

102	Optimization of aptamer microarray technology for multiple protein targets. <i>Analytica Chimica Acta</i> , 2006 , 564, 82-90	6.6	150
101	Aptamer therapeutics advance. Current Opinion in Chemical Biology, 2006, 10, 282-9	9.7	314
100	Systematic profiling of cellular phenotypes and gene function using spotted cellular microarrays. <i>FASEB Journal</i> , 2006 , 20, LB61	0.9	
99	Systematic profiling of cellular phenotypes with spotted cell microarrays reveals new mating pheromone response genes. <i>FASEB Journal</i> , 2006 , 20, A928	0.9	1
98	Binding of herpes simplex virus-1 US11 to specific RNA sequences. <i>Nucleic Acids Research</i> , 2005 , 33, 609	ን ውፊ ሷ 0	28
97	Evolution. Changing the cofactor diet of an enzyme. <i>Science</i> , 2005 , 310, 454-5	33.3	7
96	Using a deoxyribozyme ligase and rolling circle amplification to detect a non-nucleic acid analyte, ATP. <i>Journal of the American Chemical Society</i> , 2005 , 127, 2022-3	16.4	174
95	Production and processing of aptamer microarrays. <i>Methods</i> , 2005 , 37, 4-15	4.6	112
94	Prospects for the De Novo Design of Nucleic Acid Biosensors 2005 , 5-43		1
93	Synthetic biology: engineering Escherichia coli to see light. <i>Nature</i> , 2005 , 438, 441-2	50.4	467
93	Synthetic biology: engineering Escherichia coli to see light. <i>Nature</i> , 2005 , 438, 441-2 Quantum-dot aptamer beacons for the detection of proteins. <i>ChemBioChem</i> , 2005 , 6, 2163-6	50.4 3.8	231
92	Quantum-dot aptamer beacons for the detection of proteins. <i>ChemBioChem</i> , 2005 , 6, 2163-6 Functional RNA microarrays for high-throughput screening of antiprotein aptamers. <i>Analytical</i>	3.8	231
92 91	Quantum-dot aptamer beacons for the detection of proteins. <i>ChemBioChem</i> , 2005 , 6, 2163-6 Functional RNA microarrays for high-throughput screening of antiprotein aptamers. <i>Analytical Biochemistry</i> , 2005 , 338, 113-23	3.8 3.1 5.8	231
92 91 90	Quantum-dot aptamer beacons for the detection of proteins. <i>ChemBioChem</i> , 2005 , 6, 2163-6 Functional RNA microarrays for high-throughput screening of antiprotein aptamers. <i>Analytical Biochemistry</i> , 2005 , 338, 113-23 Arginine-rich motifs present multiple interfaces for specific binding by RNA. <i>Rna</i> , 2005 , 11, 1848-57 A modified consensus approach to mutagenesis inverts the cofactor specificity of Bacillus	3.8 3.1 5.8	231 83 94
92 91 90 89	Quantum-dot aptamer beacons for the detection of proteins. <i>ChemBioChem</i> , 2005 , 6, 2163-6 Functional RNA microarrays for high-throughput screening of antiprotein aptamers. <i>Analytical Biochemistry</i> , 2005 , 338, 113-23 Arginine-rich motifs present multiple interfaces for specific binding by RNA. <i>Rna</i> , 2005 , 11, 1848-57 A modified consensus approach to mutagenesis inverts the cofactor specificity of Bacillus stearothermophilus lactate dehydrogenase. <i>Protein Engineering, Design and Selection</i> , 2005 , 18, 369-77	3.8 3.1 5.8	231839421
92 91 90 89 88	Quantum-dot aptamer beacons for the detection of proteins. <i>ChemBioChem</i> , 2005 , 6, 2163-6 Functional RNA microarrays for high-throughput screening of antiprotein aptamers. <i>Analytical Biochemistry</i> , 2005 , 338, 113-23 Arginine-rich motifs present multiple interfaces for specific binding by RNA. <i>Rna</i> , 2005 , 11, 1848-57 A modified consensus approach to mutagenesis inverts the cofactor specificity of Bacillus stearothermophilus lactate dehydrogenase. <i>Protein Engineering, Design and Selection</i> , 2005 , 18, 369-77 Direct selection of trans-acting ligase ribozymes by in vitro compartmentalization. <i>Rna</i> , 2005 , 11, 1555-	3.8 3.1 5.8	231 83 94 21 45

84	Evolution of a T7 RNA polymerase variant that transcribes 2TO-methyl RNA. <i>Nature Biotechnology</i> , 2004 , 22, 1155-60	44.5	137
83	The importance of prebiotic chemistry in the RNA world. <i>Current Opinion in Chemical Biology</i> , 2004 , 8, 629-33	9.7	16
82	Aptamer-based sensor arrays for the detection and quantitation of proteins. <i>Analytical Chemistry</i> , 2004 , 76, 4066-75	7.8	274
81	Micromechanical detection of proteins using aptamer-based receptor molecules. <i>Analytical Chemistry</i> , 2004 , 76, 3194-8	7.8	276
80	Evolving new genetic codes. <i>Trends in Ecology and Evolution</i> , 2004 , 19, 69-75	10.9	37
79	Peptide-templated nucleic acid ligation. <i>Journal of Molecular Evolution</i> , 2003 , 56, 607-15	3.1	22
78	Evolution of phage with chemically ambiguous proteomes. <i>BMC Evolutionary Biology</i> , 2003 , 3, 24	3	31
77	Simultaneous detection of diverse analytes with an aptazyme ligase array. <i>Analytical Biochemistry</i> , 2003 , 312, 106-12	3.1	70
76	Survival and polymerase chain reaction-based detection of nucleic acid taggant markers during bacterial growth and sterilization. <i>Analytica Chimica Acta</i> , 2003 , 475, 67-73	6.6	1
75	Exponential growth by cross-catalytic cleavage of deoxyribozymogens. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2003 , 100, 6416-21	11.5	69
74	In vitro selection of molecular beacons. <i>Nucleic Acids Research</i> , 2003 , 31, 5700-13	20.1	72
73	Group I aptazymes as genetic regulatory switches. <i>BMC Biotechnology</i> , 2002 , 2, 21	3.5	74
72	Protein-dependent ribozymes report molecular interactions in real time. <i>Nature Biotechnology</i> , 2002 , 20, 717-22	44.5	144
71	Selecting nucleic acids for biosensor applications. <i>Combinatorial Chemistry and High Throughput Screening</i> , 2002 , 5, 263-70	1.3	50
70	Automated selection of aptamers against protein targets translated in vitro: from gene to aptamer. <i>Nucleic Acids Research</i> , 2002 , 30, e108	20.1	135
69	Increasing the thermal stability of an oligomeric protein, beta-glucuronidase. <i>Journal of Molecular Biology</i> , 2002 , 315, 325-37	6.5	52
68	Automated acquisition of aptamer sequences. <i>Combinatorial Chemistry and High Throughput Screening</i> , 2002 , 5, 289-99	1.3	53
67	Nucleic Acids for Reagentless Biosensors 2002 , 369-396		5

66	Automated selection of anti-protein aptamers. <i>Bioorganic and Medicinal Chemistry</i> , 2001 , 9, 2525-31	3.4	318
65	Aptamer beacons for the direct detection of proteins. <i>Analytical Biochemistry</i> , 2001 , 294, 126-31	3.1	525
64	In vitro selection of nucleoprotein enzymes. <i>Nature Biotechnology</i> , 2001 , 19, 650-5	44.5	81
63	Selection and characterization of Escherichia coli variants capable of growth on an otherwise toxic tryptophan analogue. <i>Journal of Bacteriology</i> , 2001 , 183, 5414-25	3.5	61
62	In vitro selection of RNA aptamers to a protein target by filter immobilization. <i>Current Protocols in Molecular Biology</i> , 2001 , Chapter 24, Unit 24.3	2.9	4
61	In vitro selection of RNA aptamers to a protein target by filter immobilization. <i>Current Protocols in Nucleic Acid Chemistry</i> , 2001 , Chapter 9, Unit 9.3	0.5	5
60	In vitro evolution of beta-glucuronidase into a beta-galactosidase proceeds through non-specific intermediates. <i>Journal of Molecular Biology</i> , 2001 , 305, 331-9	6.5	153
59	Optimization and optimality of a short ribozyme ligase that joins non-Watson-Crick base pairings. <i>Rna</i> , 2001 , 7, 513-23	5.8	33
58	Characteristics of amino acids. Current Protocols in Molecular Biology, 2001, Appendix 1, Appendix 1C	2.9	4
57	In vitro selection of signaling aptamers. <i>Nature Biotechnology</i> , 2000 , 18, 1293-7	44.5	249
57 56	In vitro selection of signaling aptamers. <i>Nature Biotechnology</i> , 2000 , 18, 1293-7 The scene of a frozen accident. <i>Rna</i> , 2000 , 6, 485-98	44·5 5.8	249
56	The scene of a frozen accident. <i>Rna</i> , 2000 , 6, 485-98		33
56 55	The scene of a frozen accident. <i>Rna</i> , 2000 , 6, 485-98 Design and optimization of effector-activated ribozyme ligases. <i>Nucleic Acids Research</i> , 2000 , 28, 1751-9 In vitro selection of nucleic acids for diagnostic applications. <i>Reviews in Molecular Biotechnology</i> ,		33 95
56 55 54	The scene of a frozen accident. <i>Rna</i> , 2000 , 6, 485-98 Design and optimization of effector-activated ribozyme ligases. <i>Nucleic Acids Research</i> , 2000 , 28, 1751-9 In vitro selection of nucleic acids for diagnostic applications. <i>Reviews in Molecular Biotechnology</i> , 2000 , 74, 15-25 Photoactivated DNA cleavage via charge transfer promoted N2 release from	920.1	339578
56 55 54 53	The scene of a frozen accident. <i>Rna</i> , 2000 , 6, 485-98 Design and optimization of effector-activated ribozyme ligases. <i>Nucleic Acids Research</i> , 2000 , 28, 1751-9 In vitro selection of nucleic acids for diagnostic applications. <i>Reviews in Molecular Biotechnology</i> , 2000 , 74, 15-25 Photoactivated DNA cleavage via charge transfer promoted N2 release from tris[3-hydroxy-1,2,3-benzotriazine-4(3H)-one]iron(III). <i>Chemical Communications</i> , 2000 , 69-70 Designed Signaling Aptamers that Transduce Molecular Recognition to Changes in Fluorescence	920.1 5.8	33957832
56 55 54 53 52	The scene of a frozen accident. <i>Rna</i> , 2000 , 6, 485-98 Design and optimization of effector-activated ribozyme ligases. <i>Nucleic Acids Research</i> , 2000 , 28, 1751-9 In vitro selection of nucleic acids for diagnostic applications. <i>Reviews in Molecular Biotechnology</i> , 2000 , 74, 15-25 Photoactivated DNA cleavage via charge transfer promoted N2 release from tris[3-hydroxy-1,2,3-benzotriazine-4(3H)-one]iron(III). <i>Chemical Communications</i> , 2000 , 69-70 Designed Signaling Aptamers that Transduce Molecular Recognition to Changes in Fluorescence Intensity. <i>Journal of the American Chemical Society</i> , 2000 , 122, 2469-2473	920.1 5.8 16.4	33 95 78 32 256

48	In vitro selection of an allosteric ribozyme that transduces analytes to amplicons. <i>Nature Biotechnology</i> , 1999 , 17, 62-6	44.5	226
47	The fidelity of template-directed oligonucleotide ligation and the inevitability of polymerase function. <i>Origins of Life and Evolution of Biospheres</i> , 1999 , 29, 375-90	1.5	34
46	Monitoring the growth of a bacteria culture by MALDI-MS of whole cells. <i>Analytical Chemistry</i> , 1999 , 71, 1990-6	7.8	143
45	Molecular parasites that evolve longer genomes. <i>Journal of Molecular Evolution</i> , 1999 , 49, 656-63	3.1	13
44	Anchoring an extended HTLV-1 Rex peptide within an RNA major groove containing junctional base triples. <i>Structure</i> , 1999 , 7, 1461-72	5.2	36
43	NMR mapping of the recombinant mouse major urinary protein I binding site occupied by the pheromone 2-sec-butyl-4,5-dihydrothiazole. <i>Biochemistry</i> , 1999 , 38, 9850-61	3.2	54
42	Polyvalent Rev decoys act as artificial Rev-responsive elements. <i>Journal of Virology</i> , 1999 , 73, 4341-9	6.6	17
41	Anti-Rex aptamers as mimics of the Rex-binding element. <i>Journal of Virology</i> , 1999 , 73, 4962-71	6.6	28
40	The limits of specificity: an experimental analysis with RNA aptamers to MS2 coat protein variants. <i>Molecular Diversity</i> , 1998 , 4, 75-89	3.1	52
39	Automated RNA selection. <i>Biotechnology Progress</i> , 1998 , 14, 845-50	2.8	117
38	Crystal structure of an RNA aptamer-protein complex at 2.8 A resolution. <i>Nature Structural Biology</i> , 1998 , 5, 133-9		118
37	Adapting selected nucleic acid ligands (aptamers) to biosensors. <i>Analytical Chemistry</i> , 1998 , 70, 3419-25	5 7.8	320
36	RNA molecules that bind to and inhibit the active site of a tyrosine phosphatase. <i>Journal of Biological Chemistry</i> , 1998 , 273, 14309-14	5.4	45
35	Incorporation of 5?-N-BOC-2?, 5?-Dideoxynucleoside-3?-O-Phosphoramidites Into Oligonucleotides by Automated Synthesis. <i>Nucleosides & Nucleotides</i> , 1997 , 16, 1821-1836		5
34	Disulfide-Intact and -Reduced Lysozyme in the Gas Phase: Conformations and Pathways of Folding and Unfolding. <i>Journal of Physical Chemistry B</i> , 1997 , 101, 3891-3900	3.4	216
33	Gas-Phase DNA: Oligothymidine Ion Conformers. <i>Journal of the American Chemical Society</i> , 1997 , 119, 9051-9052	16.4	76
32	Nucleic Acid Selection and the Challenge of Combinatorial Chemistry. <i>Chemical Reviews</i> , 1997 , 97, 349-3	3 78 .1	453
31	Cofactor-Assisted Self-Cleavage in DNA Libraries with a 3?\(\bar{B}\)?-Phosphoramidate Bond. <i>Angewandte Chemie International Edition in English</i> , 1997 , 36, 1321-1324		46

30	A biopolymer by any other name would bind as well: a comparison of the ligand-binding pockets of nucleic acids and proteins. <i>Structure</i> , 1997 , 5, 729-34	5.2	22
29	Surprising fidelity of template-directed chemical ligation of oligonucleotides. <i>Chemistry and Biology</i> , 1997 , 4, 595-605		39
28	Screening chemical libraries for nucleic-acid-binding drugs by in vitro selection: a test case with lividomycin. <i>Molecular Diversity</i> , 1996 , 2, 103-10	3.1	14
27	Effects of Matrix Variations and the Presence of Iron on Matrix-assisted Laser Desorption/Ionization Mass Spectra of DNA. <i>Rapid Communications in Mass Spectrometry</i> , 1996 , 10, 198	30 2 1386	5 ¹⁴
26	Deep penetration of an alpha-helix into a widened RNA major groove in the HIV-1 rev peptide-RNA aptamer complex. <i>Nature Structural Biology</i> , 1996 , 3, 1026-33		146
25	DNA shuffling brightens prospects for GFP. <i>Nature Biotechnology</i> , 1996 , 14, 366	44.5	7
24	The search for missing links between self-replicating nucleic acids and the RNA world. <i>Origins of Life and Evolution of Biospheres</i> , 1995 , 25, 515-30	1.5	31
23	High resolution matrix-assisted laser desorption/ionization time-of-flight analysis of single-stranded DNA of 27 to 68 nucleotides in length. <i>Rapid Communications in Mass Spectrometry</i> , 1995 , 9, 1061-6	2.2	42
22	In vitro selection of RNA lectins: using combinatorial chemistry to interpret ribozyme evolution. <i>Chemistry and Biology</i> , 1995 , 2, 291-303		92
21	Artificial evolution and natural ribozymes. <i>FASEB Journal</i> , 1995 , 9, 1183-95	0.9	36
20	Re-creating the RNA world. Current Biology, 1995, 5, 1017-22	6.3	39
19	Using in vitro nucleic acid selections for conventional drug design. <i>Drug Development Research</i> , 1994 , 33, 102-115	5.1	9
18	A three-dimensional model of the Rev-binding element of HIV-1 derived from analyses of aptamers. <i>Nature Structural and Molecular Biology</i> , 1994 , 1, 293-300	17.6	48
17	RNA selection. Aptamers achieve the desired recognition. <i>Current Biology</i> , 1994 , 4, 427-9	6.3	92
16	The RNA world. Empirical explorations of sequence space: Host-guest chemistry in the RNA world. <i>Zeitschrift Fur Elektrotechnik Und Elektrochemie</i> , 1994 , 98, 1115-1121		13
15	Selection and design of high-affinity RNA ligands for HIV-1 Rev. <i>Gene</i> , 1993 , 137, 19-24	3.8	55
14	Selective optimization of the Rev-binding element of HIV-1. <i>Nucleic Acids Research</i> , 1993 , 21, 5509-16	20.1	127
13	Selection in vitro of single-stranded DNA molecules that fold into specific ligand-binding structures. <i>Nature</i> , 1992 , 355, 850-2	50.4	643

12	In vitro selection of RNA molecules that bind specific ligands. <i>Nature</i> , 1990 , 346, 818-22	50.4	7326
11	In vitro genetic analysis of the Tetrahymena self-splicing intron. <i>Nature</i> , 1990 , 347, 406-8	50.4	120
10	Phylogenetic and genetic evidence for base-triples in the catalytic domain of group I introns. <i>Nature</i> , 1990 , 347, 578-80	50.4	133
9	STANDARD MEASUREMENTS, DATA, AND ABBREVIATIONS. <i>Current Protocols in Molecular Biology</i> , 1990 , 12, A.1.1-A.1.32	2.9	
8	Purification of Oligonucleotides Using Denaturing Polyacrylamide Gel Electrophoresis: Preparation and Analysis of DNA. <i>Current Protocols in Molecular Biology</i> , 1989 , 6, 2.12.1-2.12.5	2.9	
7	Biocompatible Materials Enabled by Biobased Production of Pyomelanin Isoforms Using an Engineered Yarrowia lipolytica. <i>Advanced Functional Materials</i> ,2109366	15.6	1
6	Deep learning redesign of PETase for practical PET degrading applications		1
5	Deep learning redesign of PETase for practical PET degrading applications High-surety isothermal amplification and detection of SARS-CoV-2, including with crude enzymes		33
5	High-surety isothermal amplification and detection of SARS-CoV-2, including with crude enzymes An improved and readily available version of Bst DNA Polymerase for LAMP, and applications to		33
5	High-surety isothermal amplification and detection of SARS-CoV-2, including with crude enzymes An improved and readily available version of Bst DNA Polymerase for LAMP, and applications to COVID-19 diagnostics		7